



Creating a public resource for protein allergenicity assessment: The COMPARE Database

November 9, 2017

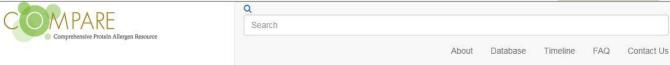


HESI COMPARE Allergen Database: www.comparedatabase.org





- A comprehensive and up to date collection of peer reviewed allergen protein sequences. Builds on a > 10 year database of peer review science.
- Includes known allergens from 2016 and previous years (listed in existing public database AllergenOnline V16) and new allergens identified through the COMPARE process.
- Available since February 03, 2017 at <u>www.comparedatabase.org</u> (website hosted and maintained by HESI)
- Updated annually



Compare Database

[Search Database | COMPARE 2017 FASTA sequences (PDF) | COMPARE 2017 FASTA sequences (Text)]

All sequence records originated from the National Center for Biotechnology Information (NCBI) database. As of 02 February 2017, all sequence records are valid

| 🛓 Download Data | | | | | | |
|---------------------|-------------|-----------|------------|--------|--------------|-----------------------|
| SPECIES A | COMMON NAME | GI# | ACCESSION# | LENGTH | YEAR ADOPTED | |
| Acarus siro | Mite | 118638268 | ABL09307.1 | 131 | 2009 | View Details |
| Acarus siro | Mite | 118638278 | ABL09312.1 | 517 | 2009 | View Details |
| Actinidia arguta | Hardy Kiwi | 441482362 | AGC39172.1 | 213 | 2014 | View Details |
| Actinidia arguta | Hardy Kiwi | 441482364 | AGC39173.1 | 213 | 2014 | View Details |
| Actinidia arguta | Hardy Kiwi | 441482366 | AGC39174.1 | 213 | 2014 | View Details |
| Actinidia chinensis | Kiwi | 68064399 | P83958.1 | 20 | 2007 | View Details |
| Actinidia chinensis | Kiwi | 190358935 | P00785.4 | 380 | 2009 | View Details |
| Actinidia chinensis | Kiwi | 281552896 | CAM31908.1 | 159 | 2011 | ₩ View Details |
| Actinidia chinensis | Kiwi | 378548410 | P85204.1 | 15 | 2013 | View Details |
| Actinidia chinensis | Kiwi | 441482354 | AGC39168.1 | 213 | 2014 | ₩ View Details |
| Actinidia chinensis | Kiwi | 441482370 | AGC39176.1 | 225 | 2014 | View Details |
| Actinidia deliciosa | Kiwi | 100 | | 109 | 2016 | View Details |
| Actinidia deliciosa | Kiwi | 15984 | CAA34486.1 | 380 | 2007 | View Details |
| Actinidia deliciosa | Kiwi | 166317 | AAA32629.1 | 380 | 2007 | ₩ View Details |
| Actinidia deliciosa | Kiwi | 40807635 | AAR92223.1 | 116 | 2007 | View Details |



Rationale for building COMPARE @ HESI:

1) Key components of the HESI public-private partnership.



HESI PATC COMPARE

P-P Science Advisory Team
Project Management Expertise

✓ Multiple stakeholders engagement.



✓ Independent coordinating body (scientific program management).

✓ Professional bioinformatics services

Peer Review Panel (PRP)

Experts

Allergy Science Expertise

✓ Decisions based on peer-review

JIFSAN and AccuraScience

Bioinformatics and Data Management Expertise



✓ Rigorous process for data management and tracking (custom information management tool).



GenBank

Rationale for building COMPARE @ HESI:

2) Good Coordination allows multiple sourcing of allergen candidates





Bioinformatics automated filtering process

IUIS

Allergen Candidate List

Literature Dbases

Targeted search strategy

Goal: enhance confidence (no allergens left behind)



Add to existing database:

COMPARE Update



Rationale for building COMPARE @ HESI: 3) Enhancing transparency with detailed documentation



What has been done:

- COMPARE overview <u>brochure</u>: at initiation of the project (Q1 2016).
- Detailed <u>process development</u> described in the COMPARE website: within 6 months of the database public release (COMPARE' 2017). http://comparedatabase.org/process-development/
- Announcement letter sent to stakeholders and published on JIFSAN's website.

In preparation:

Peer-review publication





COMPREHENSIVE PROTEIN ALLERGEN RESOURCE

Program Overview – February 2016

ABOUT HESI

HESI, the Health and
Environmental Sciences
institute, is a non-profit,
global scientific organization
iedicated to developing
science for a safer, more
sustainable world.

HESI's mission is to collaboratively identify and help to resolve global health and environmental challenges through engagement of scientists from academia, government, industry, NGOs, and other strategic partners.



OVERVIEW

The COMPARE Database, to be made publicly available in January 2017, will be a transparent resource for identification of protein sequences that are known or putative allergens. The COMPARE database will meet needs for allergy safety assessment via an annual updating process that 1) captures new listings of allergen sequences, 2) filters out non-allergen sequences, 3) identifies published literature linked to the identified potential allergen sequences, and 4) verifies that the newly identified sequences have clinical, published evidence of allergenicity based on standardized criteria.

PROGRAM STRUCTURE

The COMPARE database is a collaborative HESI program that combines programmatic support from the Joint Institute for Food Safety and Nutrition – www.jifsan.org at the University of Maryland with financial support from private sector partners. Additional in-kind support from multiple academic and government partners is engaged throughout the process. A public-private HESI collaborative team will publicly document all database design and search algorithm decisions. An exclusively public-sector expert panel will review the output from a high-throughput bioinformatic pipeline against peer-reviewed publications. The expert panel will then select sequences for inclusion in the database and document their rationale for inclusion.

TIMELIN

The COMPARE database will be released to the public by January, 2017. To enable this timing, a steering team will convene in early 2016 to ensure timelines, process, and resources are aligned. Sequence searching will be conducted in second quarter 2016 and peer-review in third quarter 2016 before final assembly and release.



Conclusion: overall program design and components enable long term, high quality and stable support.



Advisory Team

- 20 Public/Private stakeholders.
- General program oversight, QC, user feedback.
- Expert knowledge
- No influence on decisions regarding sequence inclusion or exclusion.

Bioinformatics and Data Communication

- AccuraSciences: keyword filtering and algorithm design; sources information for database.
- University of Maryland / JIFSAN:
- Literature search and procurement.
- Data communication.
- Review Tool.



Program Coordination

- Overall program management.
- Coordination between partners, PRP assembly and management.
- Database 'home' and maintenance.



Expert Peer Review Panel

- Invited public sector allergy experts.
- Define inclusion /exclusion criteria.
- Review of sequences and references / publications.
- Provide final decisions on new sequences to include in database

Participating Organizations

- Public Sector:
- Universities of Amsterdam
- University of Maryland / JIFSAN
- U.S. EPA
- U.S. FDA

Private Sector:

- BASF
- Bayer CropScience
- Dow
- DuPont Pioneer
- KWS
- Monsanto
- Syngenta
- Vilmorin & Cie







Who will use the COMPARE Database?



Who?

 Target audiences: regulatory agencies as well as the public (lay person(s) and professionals)

Rationale?

 Truly public resource: any/all stakeholders or users of the database have equal and full access. Ensures consistency and clarity.

Hows

- Sequence comparisons: these can be made between biotech proteins and known allergens – COMPARE allows these bioinformatic comparisons.
- Basic research on types of allergens, sequence information and literature associated with clinical relevance for including specific allergens.
- Staying updated with new allergens identified each year (all users).

Access to a transparent and consensus-based allergen database is a key aspect for supporting public safety.



For more information:



Visit:

- ✓ HESI PATC website: committee fact sheet, events and publications:
 http://hesiglobal.org/protein-allergenicity-technical-committee-patc/
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www.hesiglobal.org

